

09854326-051101
TOT150" 92E45860

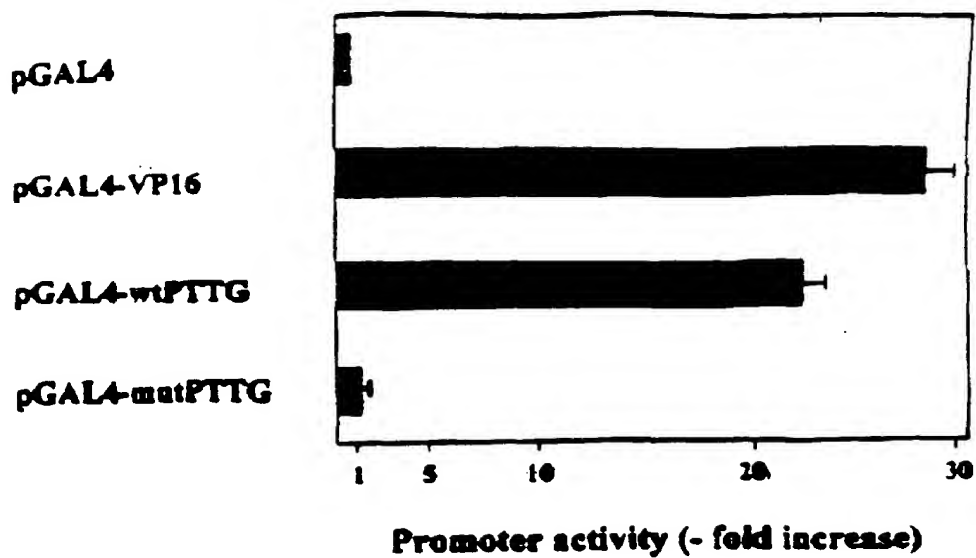


Figure 1

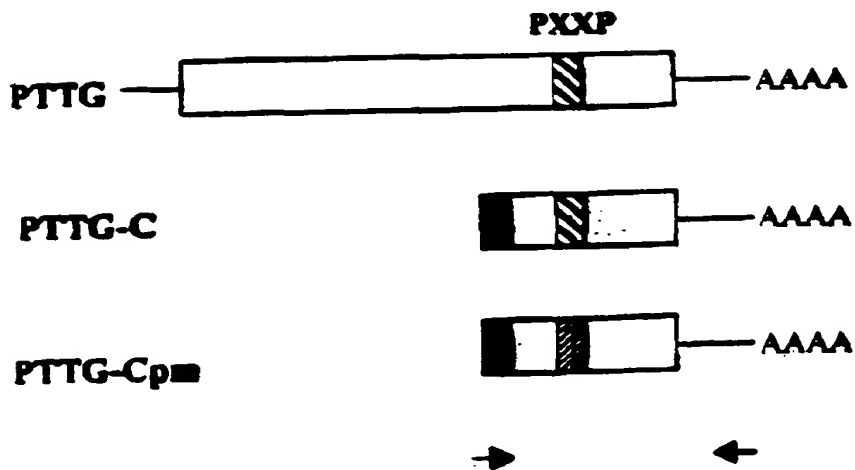


Figure 2A

09854326.054101

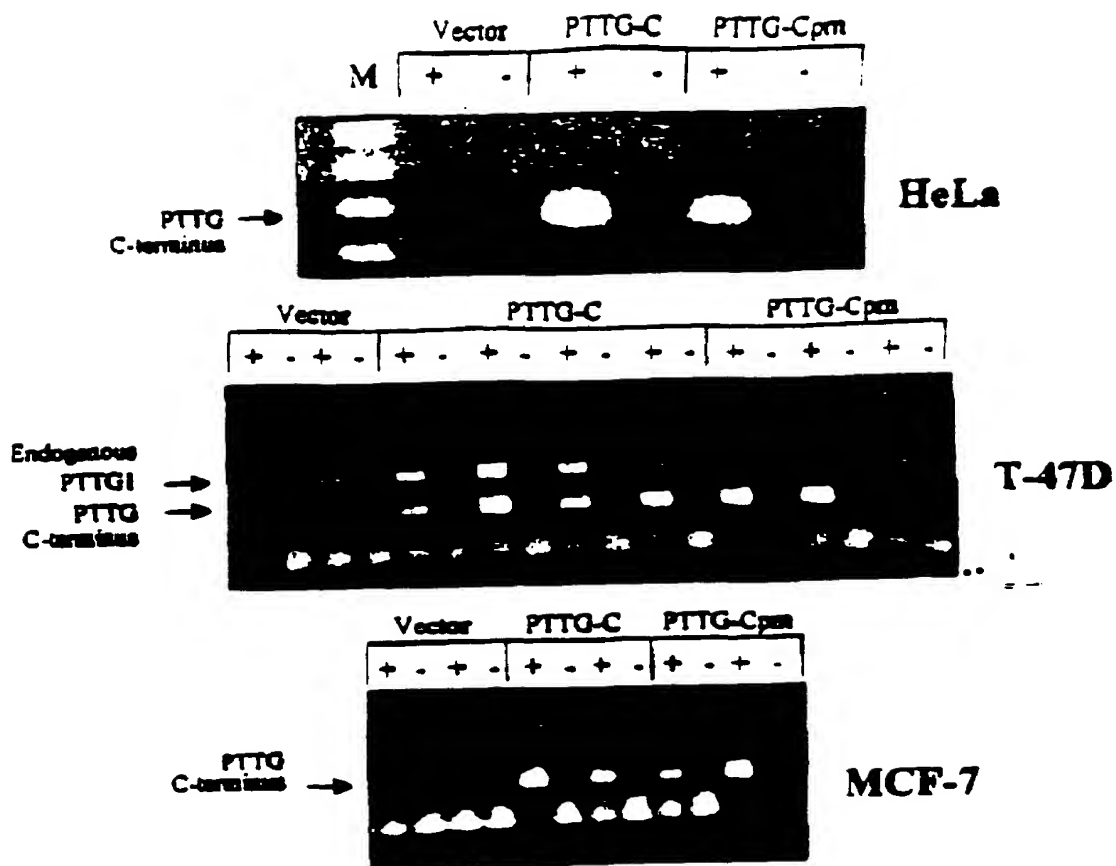


Figure 2B

5'-TTT CAG CTG GGC CCC CCT TCA CCT GTG AAG ATG CCC TCT CCA TGG GAA-3'
 P S P P P P
 A C G T A C G T
 PTTG-C PTTG-Cpm
 5'-TTT CAG CTG GGC CCC CCT CAA CTT GTG AAG ATG CTC TCC GCA CTC TGG GAA-3'
 A Q L L A L

Figure 2C

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TOTTSD" 92E45860

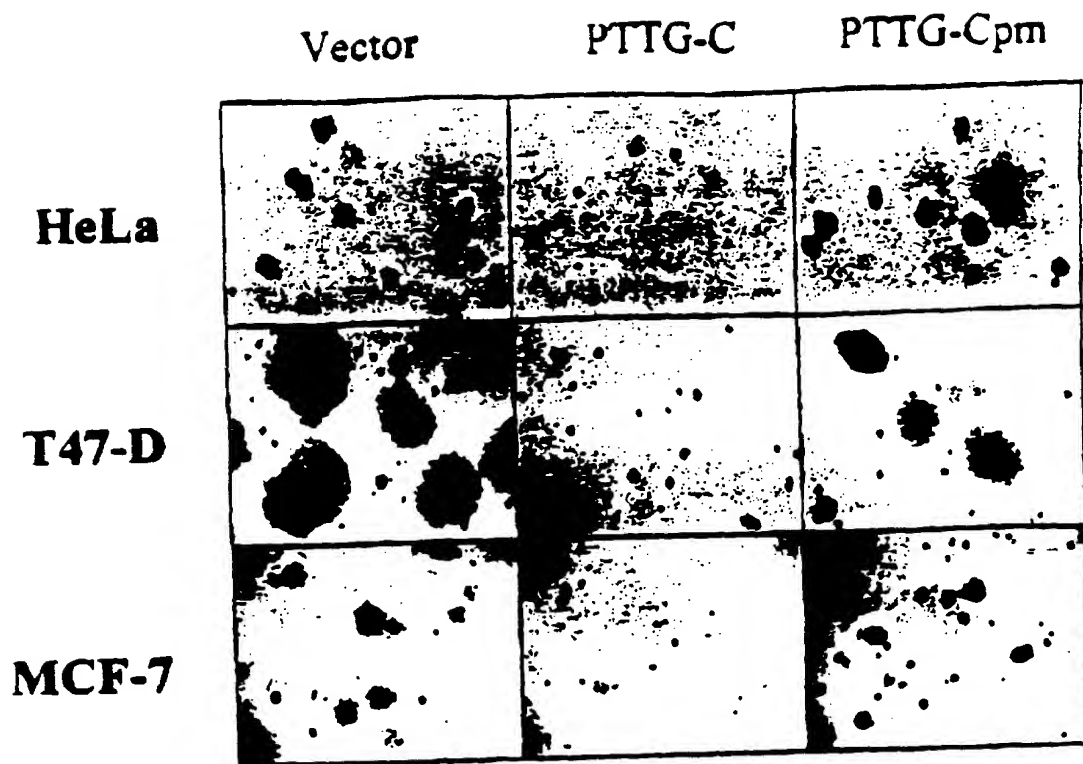


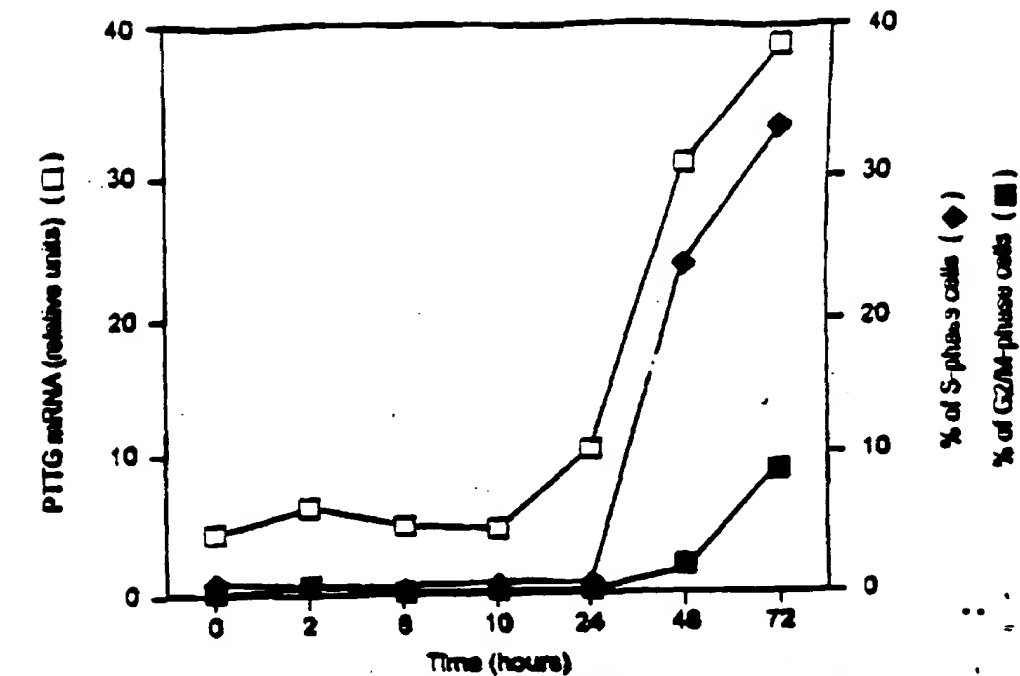
Figure 3

A bar graph showing the levels of basic FGF (pg/ml) for three groups: Vector, PTTG-C, and PTTG-Cpm. The Y-axis ranges from 0 to 4 pg/ml. The Vector group shows a high level of basic FGF (~3.0 pg/ml). The PTTG-C group shows a significantly lower level (~0.4 pg/ml). The PTTG-Cpm group shows a level intermediate between Vector and PTTG-C (~1.8 pg/ml). Error bars are present for all groups.

Group	basic FGF (pg/ml)
Vector	~3.0
PTTG-C	~0.4
PTTG-Cpm	~1.8

Figure 4.

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pttg →



28S →

18S →

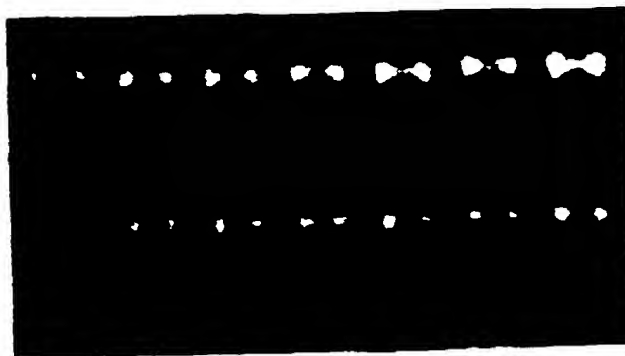


Figure 5

09854326.051101
T0TFS0" 92E45860

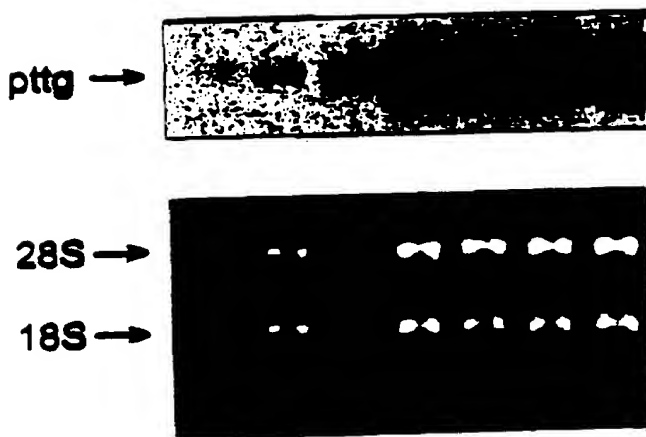
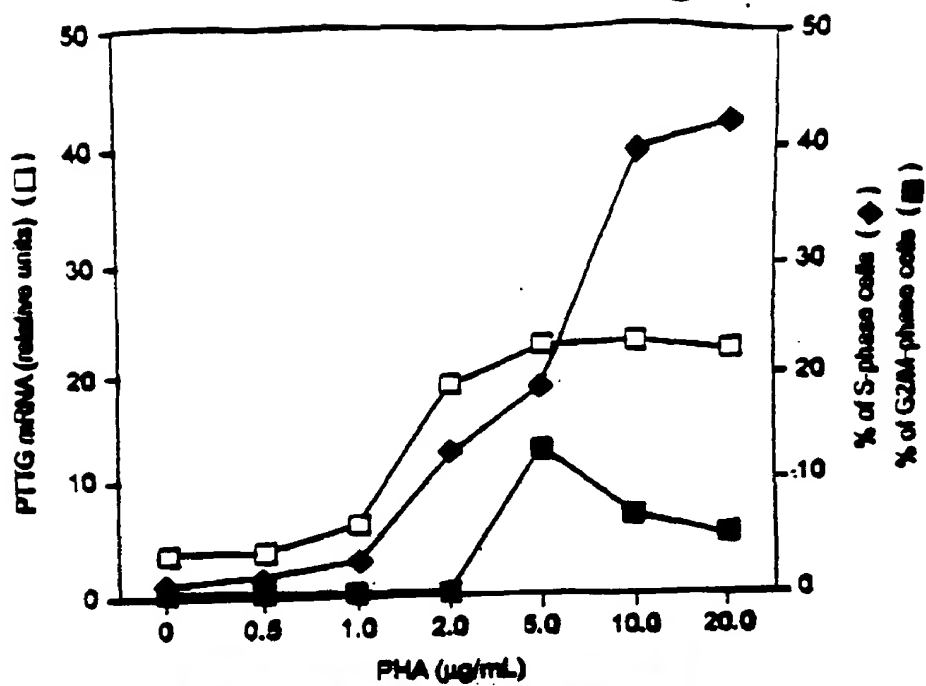


Figure 6

09854326.051103

Q3

IL-2

28S →

18S →

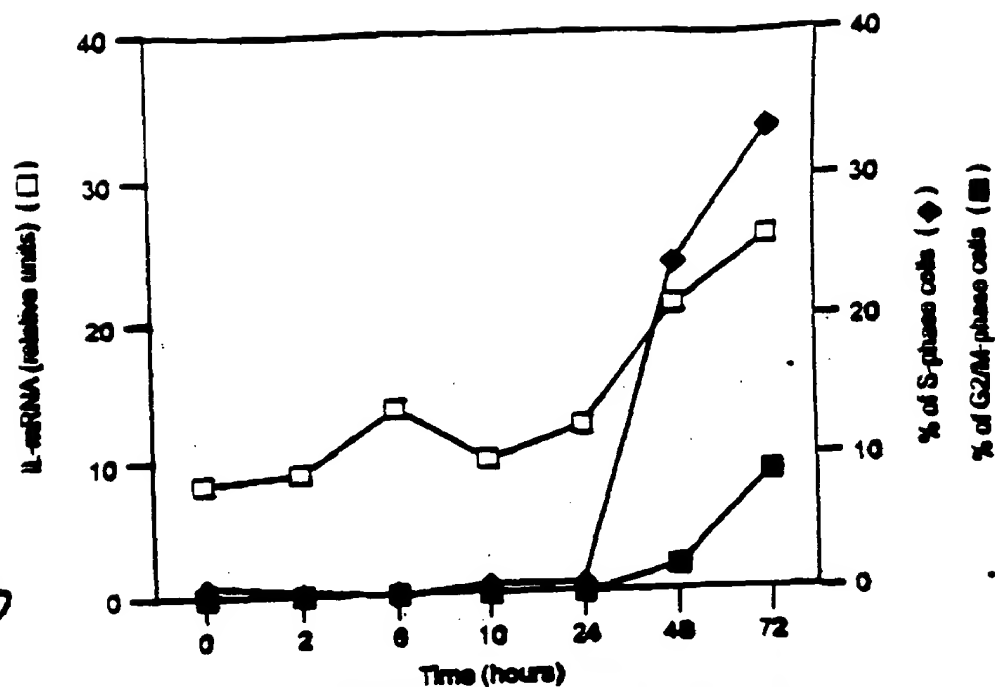


Figure 7

09854326-051101

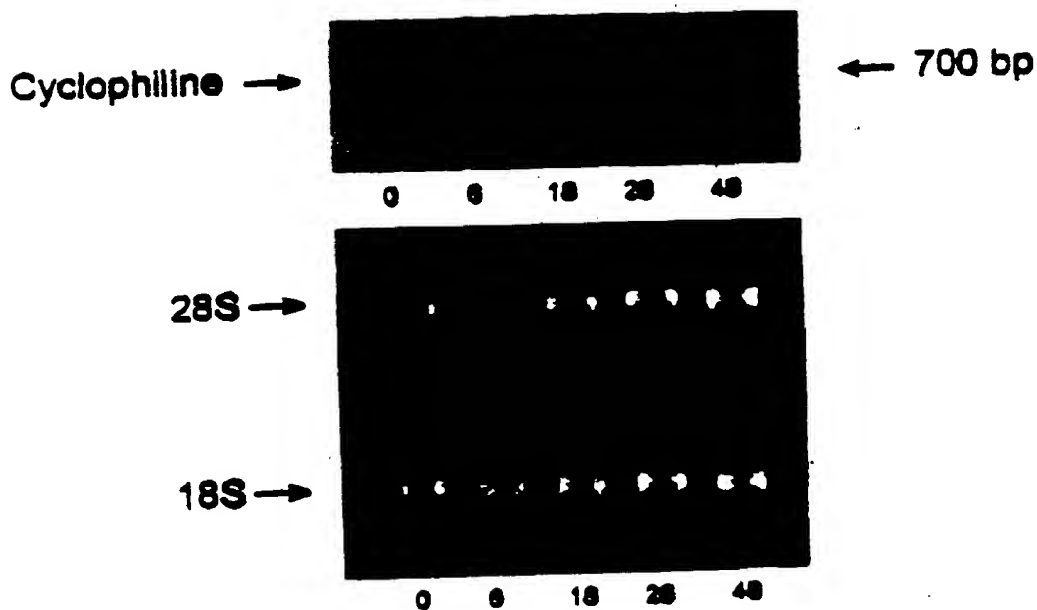
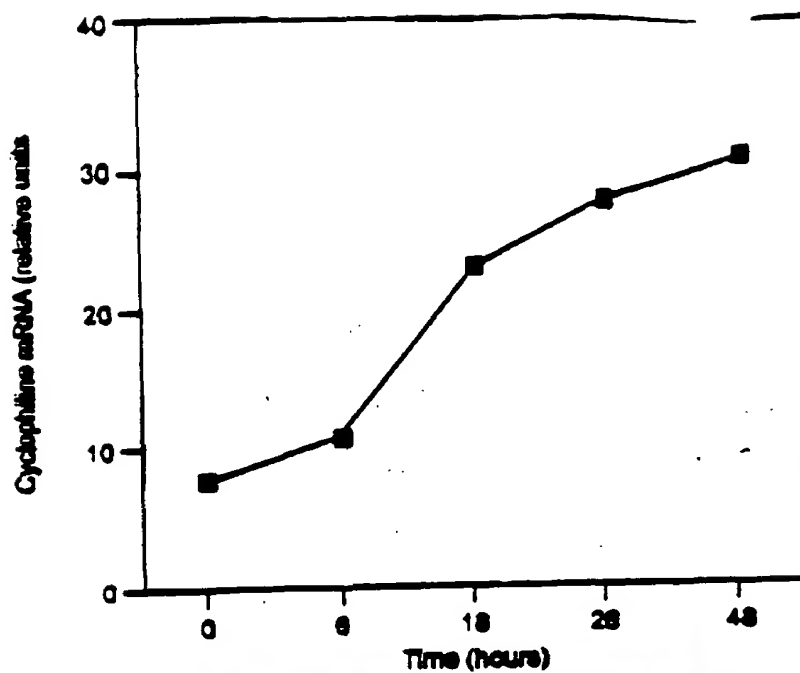


Figure 8

09854326-051101

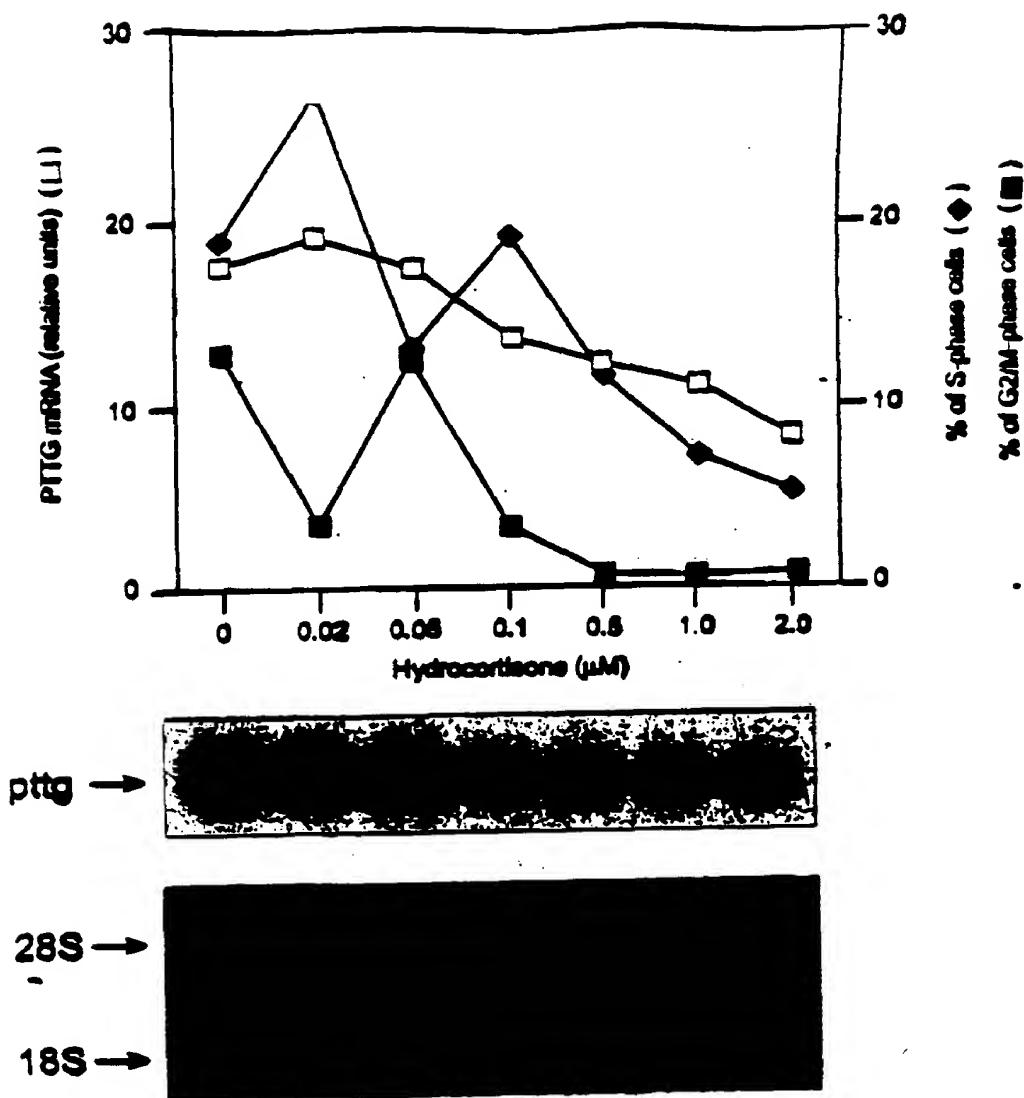


Figure 9

09254326-051101

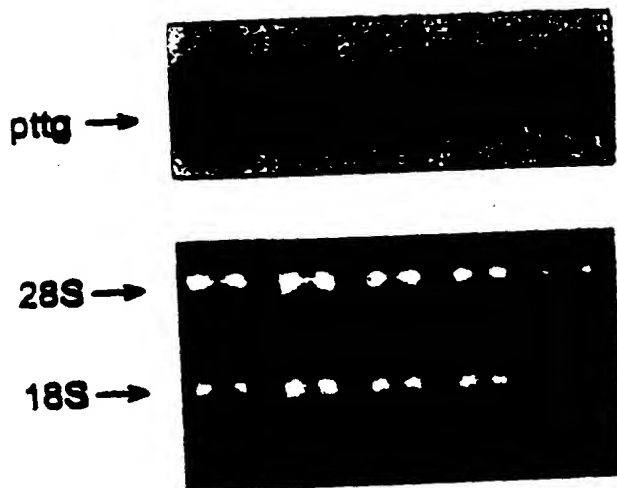
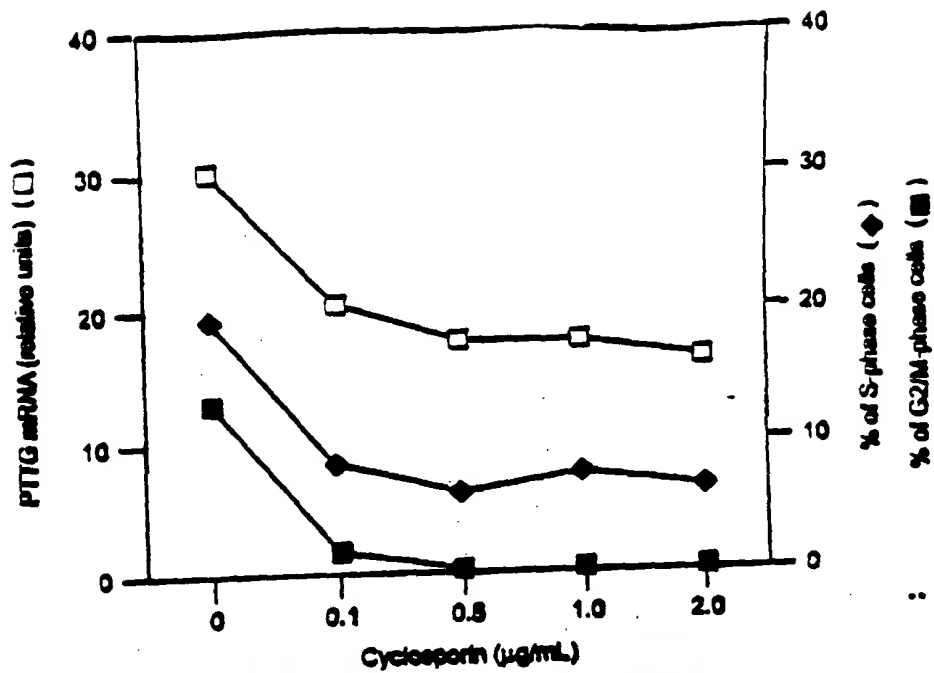
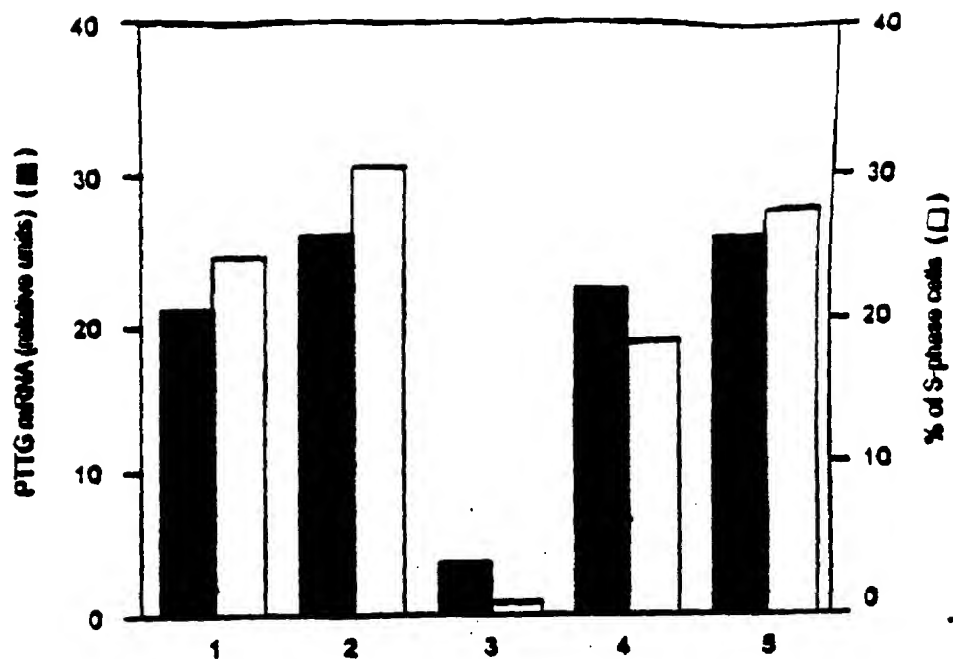


Figure 10

09854326-051101



pttg →



28S →

18S →

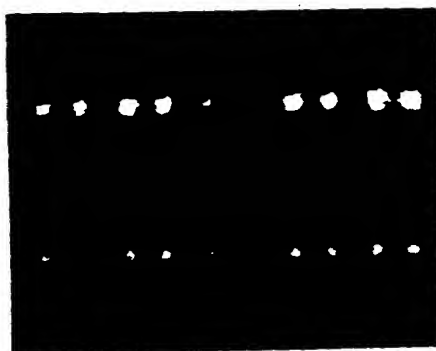


Figure 11

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TOT 50" 92E45860

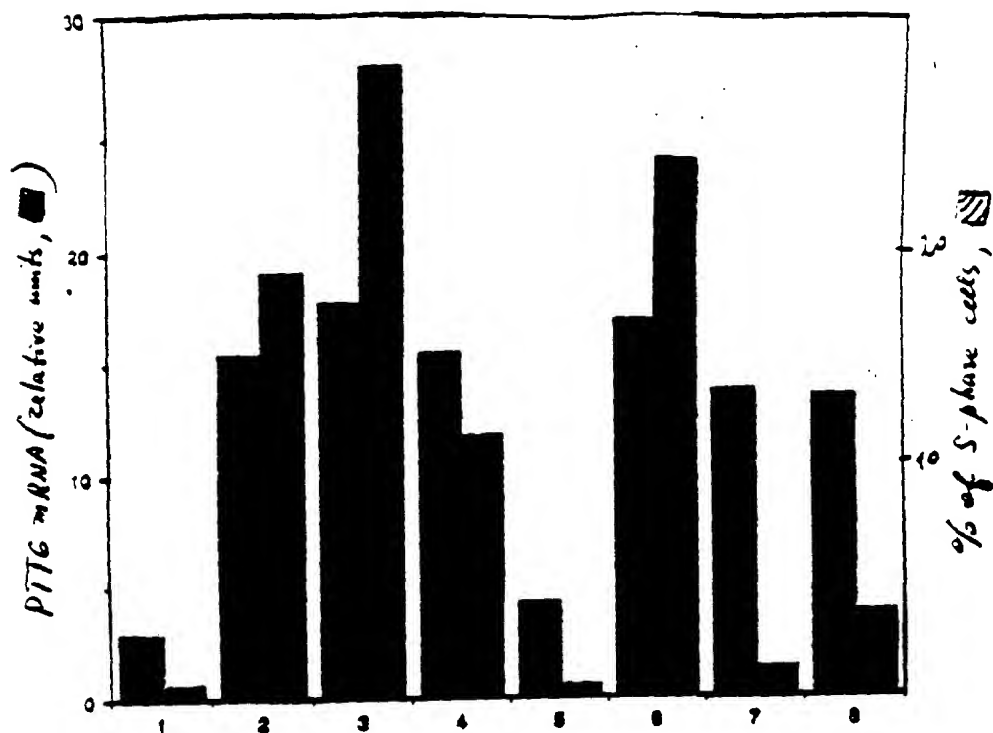


Figure 12

00854326.051104

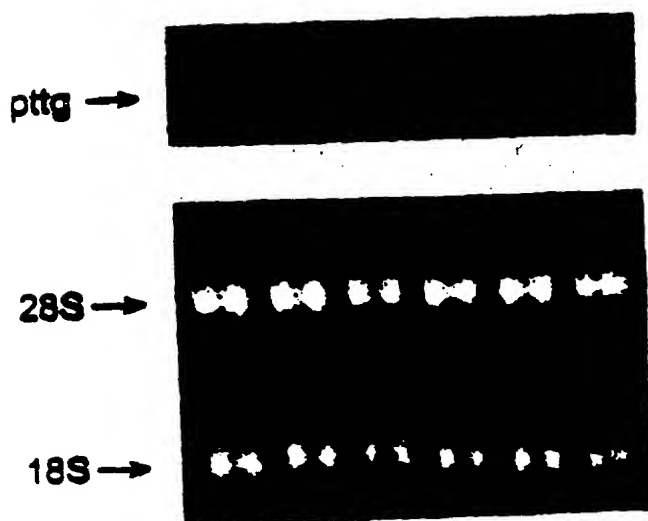
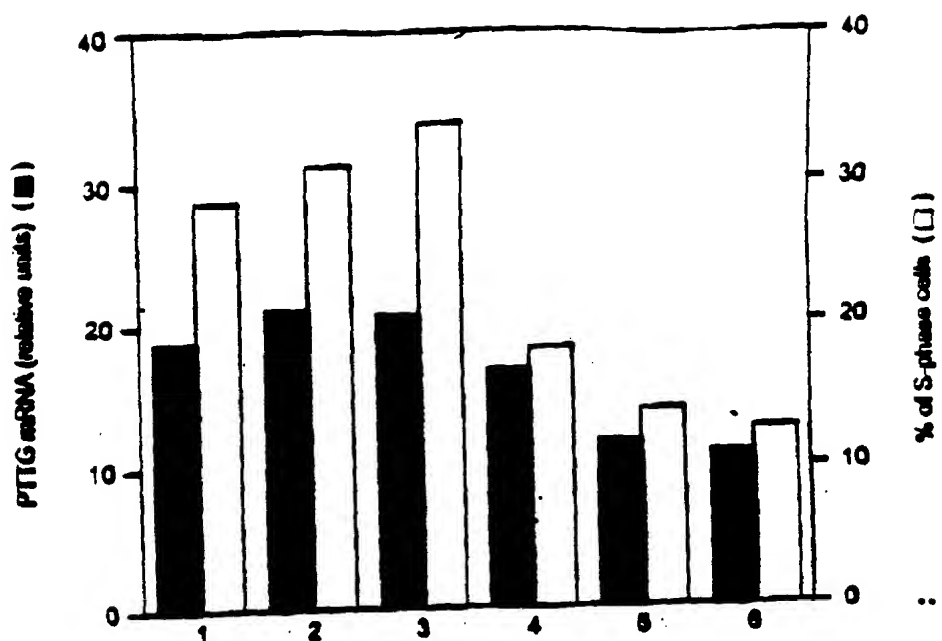


Figure 13

TOTFSD" 92E45860

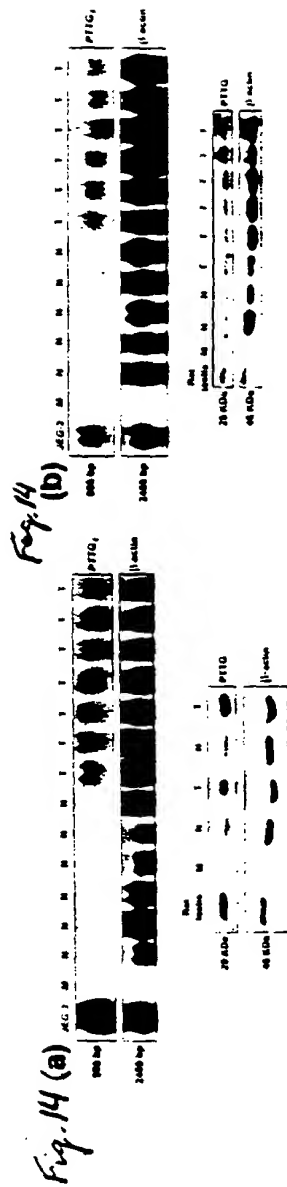


Figure 15

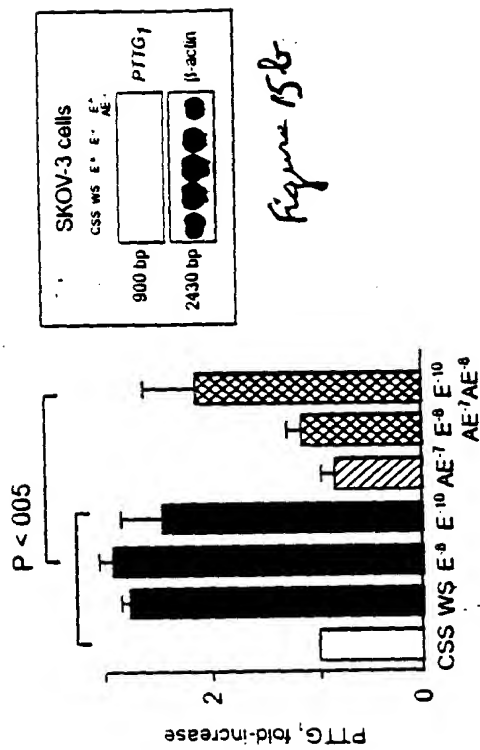


Figure 15b



Figure 15a

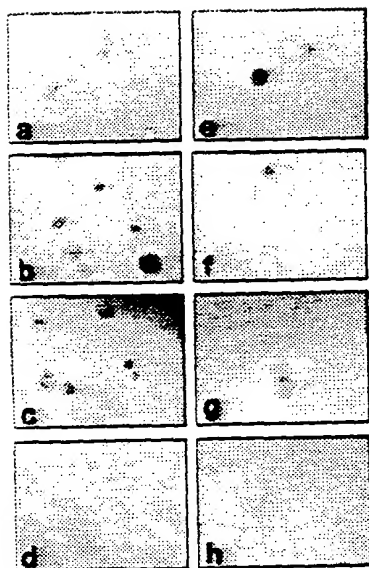


Figure 16

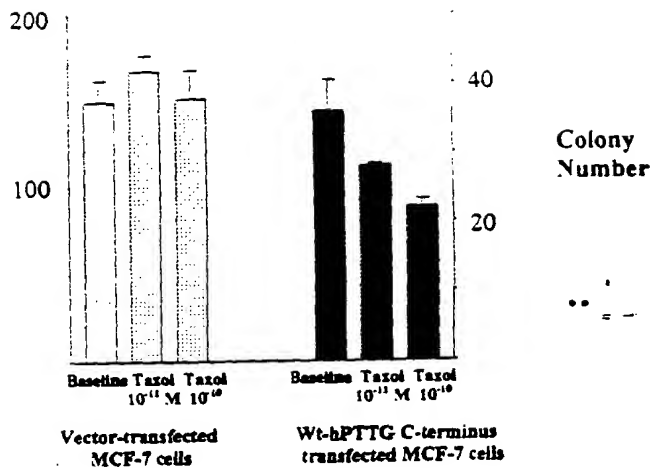


Figure 17

92E45860

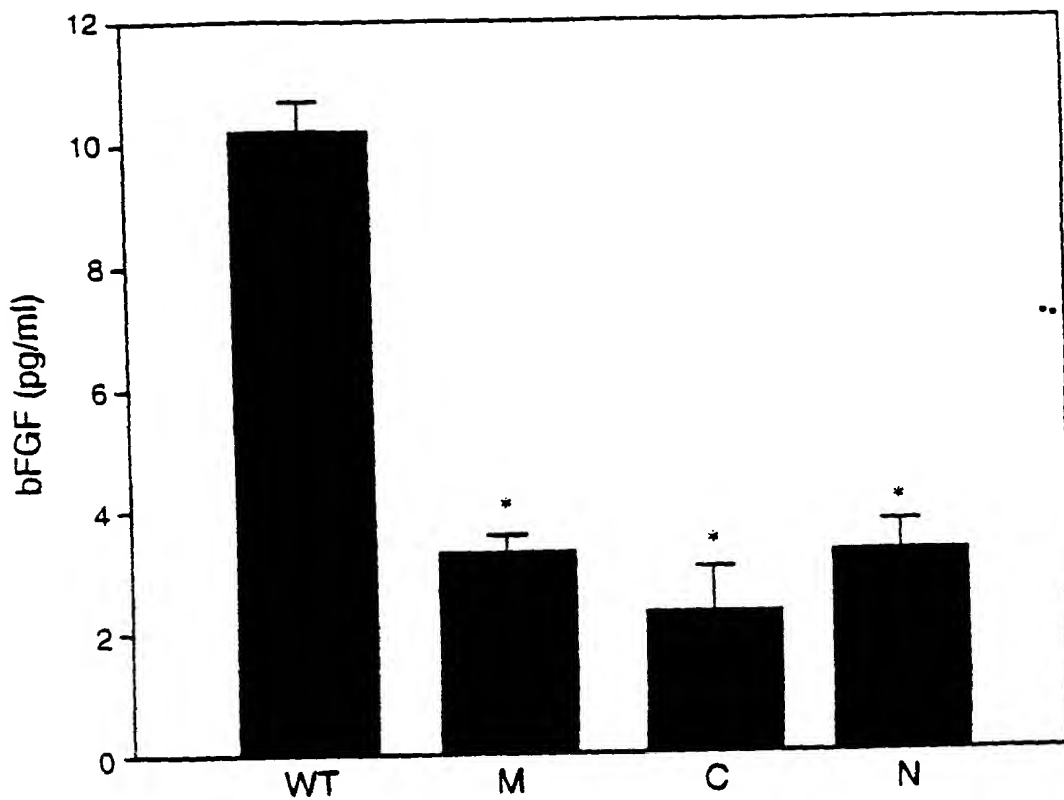


Figure 18

00854326 92E45860

Fig. 20A

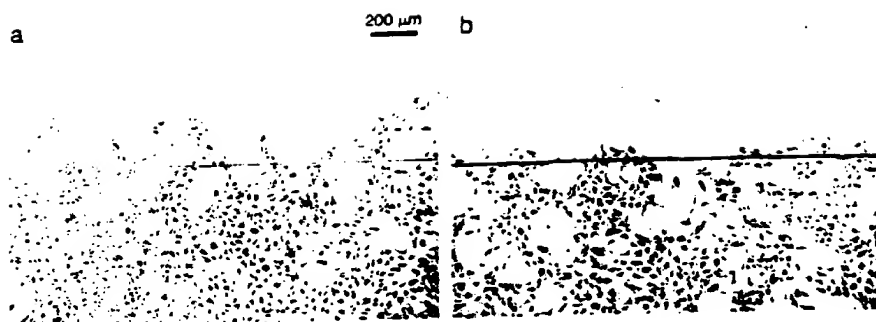


Fig 20B

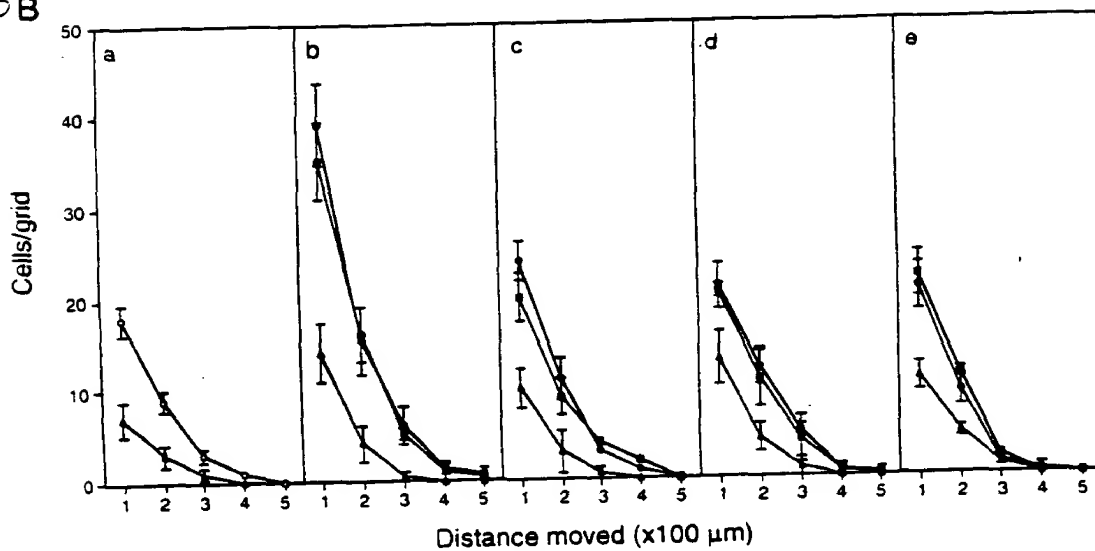


Figure 20

09854326.051101

Fig. 21A

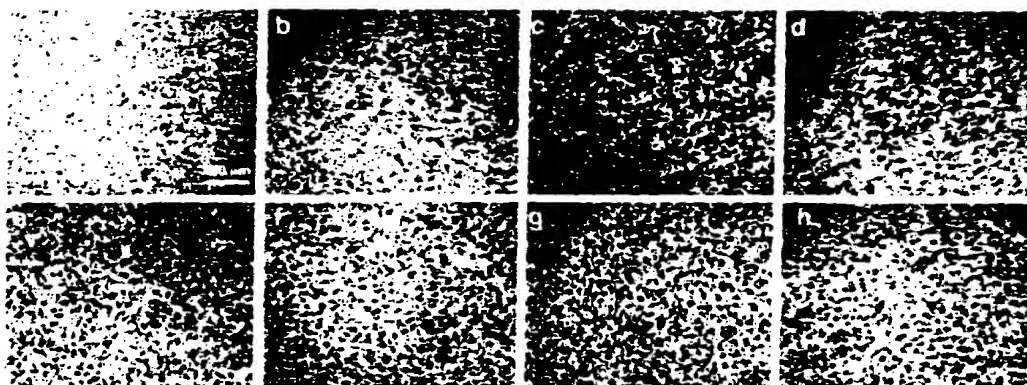


Fig. 21B

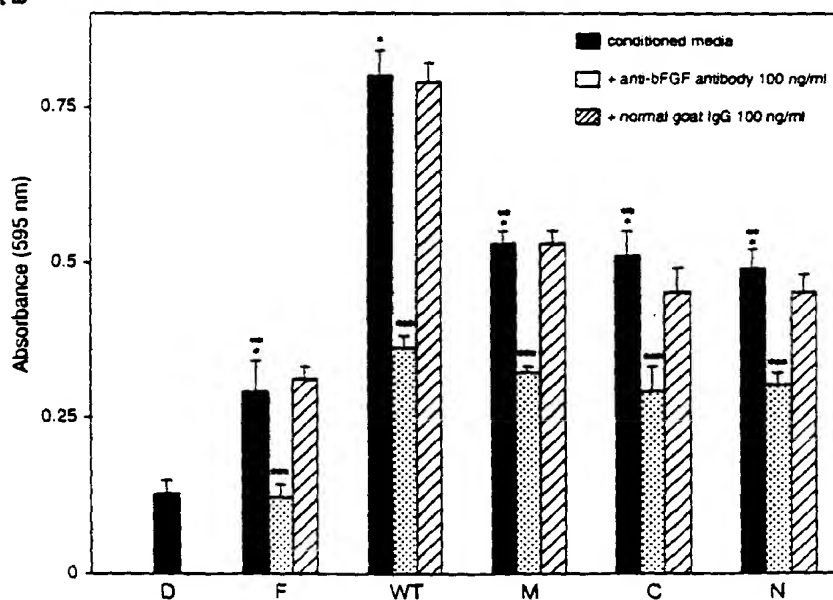


Figure 21

09854326-051101

Fig 22A

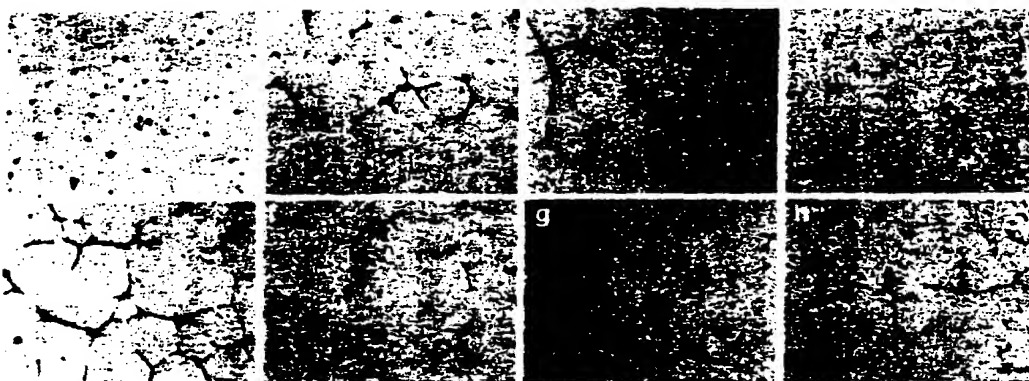


Fig 22B

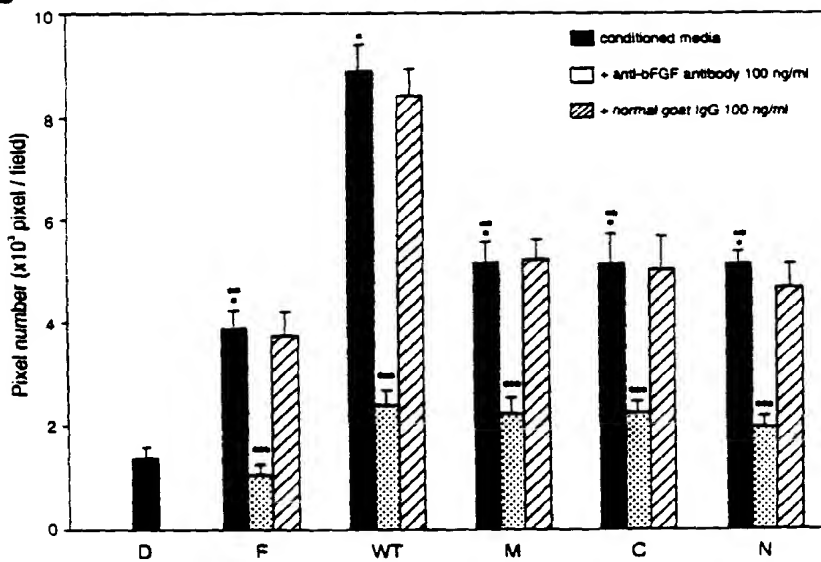
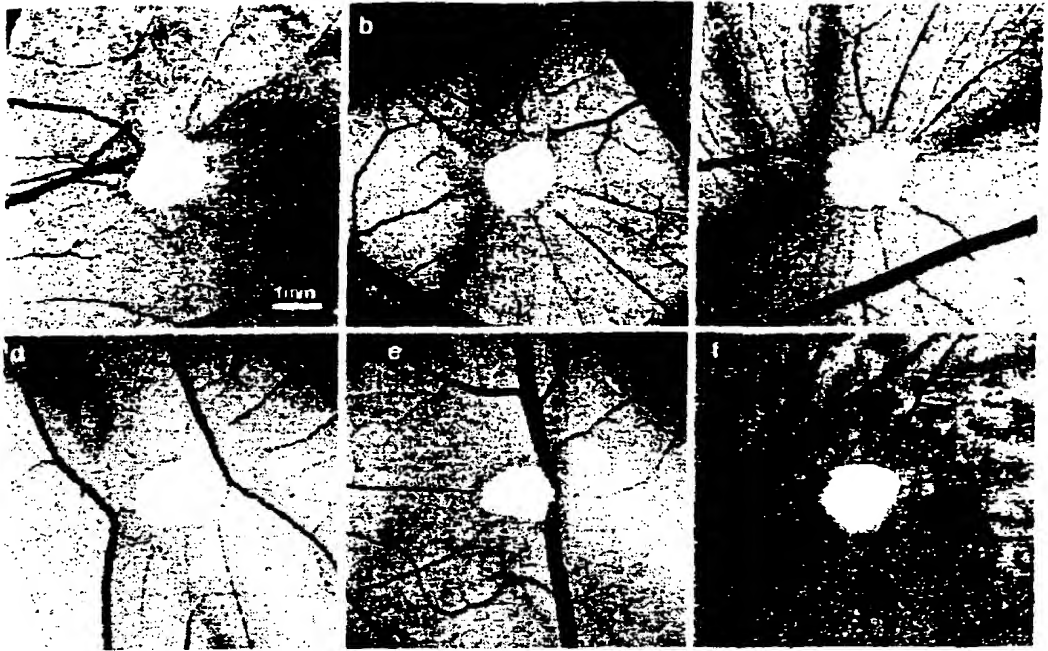


Figure 22

Fig 23A



09854326.051101

Fig. 23B

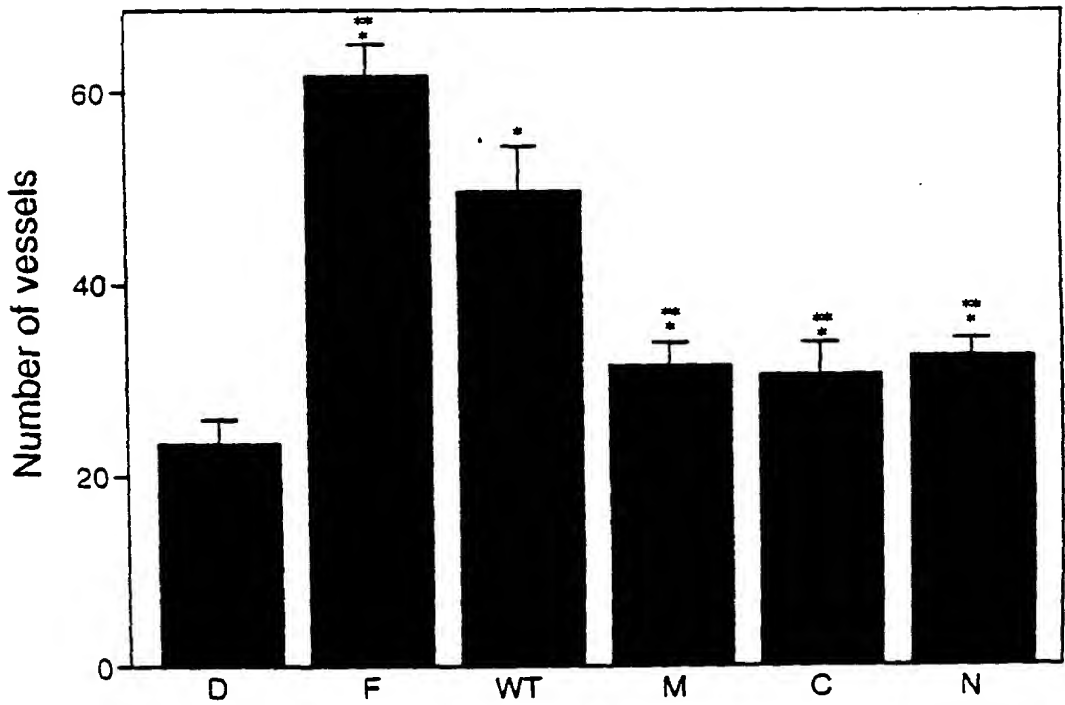


Figure 23

hPTTG1	1	MATLIYVDKE	NCEPGTRVVA	KDGLKLGSGP	SIKALDGRSQ	VSTPRFGKTF	DAPPALPKAT
hPTTG2	1	*****	I*****	**V**ER*	*****I**	*L*****Y	***S*****
hPTTG3	1	*****	*E**IL*AT	*****	*****	**ISC*****	***TS*****
mPTTG	1	*****F**D	*E**R*IAS	*****T*V	*****KL*	*****V**V*	N*-**V***S
rPTTG	1	*****F**D	*E**S*LAS	*****	*****KL*	*****V**V*	G*-*GL***S
	61	RKALGTVNRA	TEKSVKTKGP	LKQKQPSFSA	-KKMTEKTVKA	KSSVLASDDA	YPEIEKFFPF
	61	*****	*****R*	*****	*****T	****p*****	*****
	61	*****	*****	*****	*****	*N**p***G	*****L***
	58	*****V	A**PM**GK*	*QP**TLTG	**I***ST* ^T	Q***p*p***	*****
	58	*****V	***P**SSK*	*QS**TLV	*I***ST* ^T	QG*AP*p***	*****
	121	NPLDFESFDL	PEEHQIAHLP	LSGVPLMILD	EERELEKLFQ	LGPPSPVKMP	SPPWESNLLQ
	121	*L*****	***R*****	*****	**G*****	*****	***C**FA
	121	***G*****	*****	**E*****	*****	*****L***	***K*****
	118	***--***	*****SL**	*N***IT*N	***G*****LH	*****L*T*	FLS***DP*Y
	118	D*****	*****SL*	*N*****N	***G*****LH	*D*****LQK*	FL***DP*P
	181	SPSSILSTLD	VELPPVCCDI	DI			
	181	VSFKHSVDPG	C				
	181	**L***L***	*****S*	**			
	175	**P*A*****	*****Y*A	**			
	178	**P*A**A**	*****Y*A	**			

(SEA. 1D. NO. 4)
 (SEA. 1D. NO. 64)
 (SEA. 1D. NO. 67)
 (SEA. 1D. NO. 14)
 (SEA. 1D. NO. 2)

Figure 24

:

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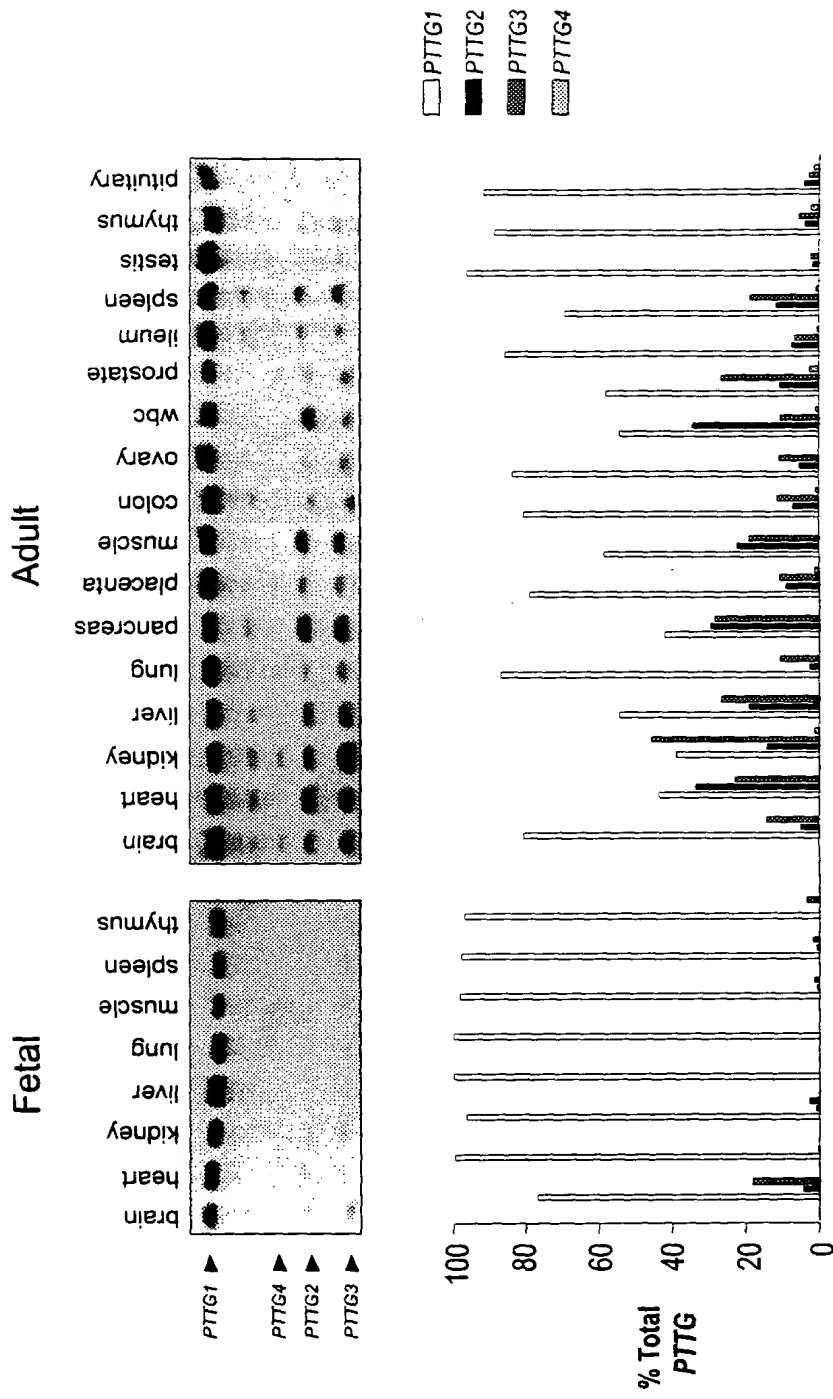


Figure 26

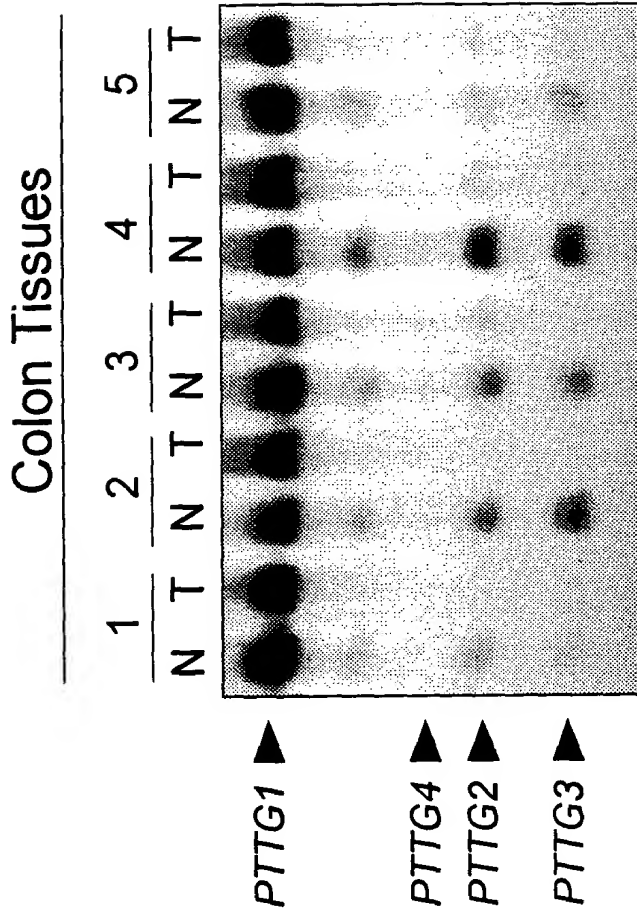


Figure 27A

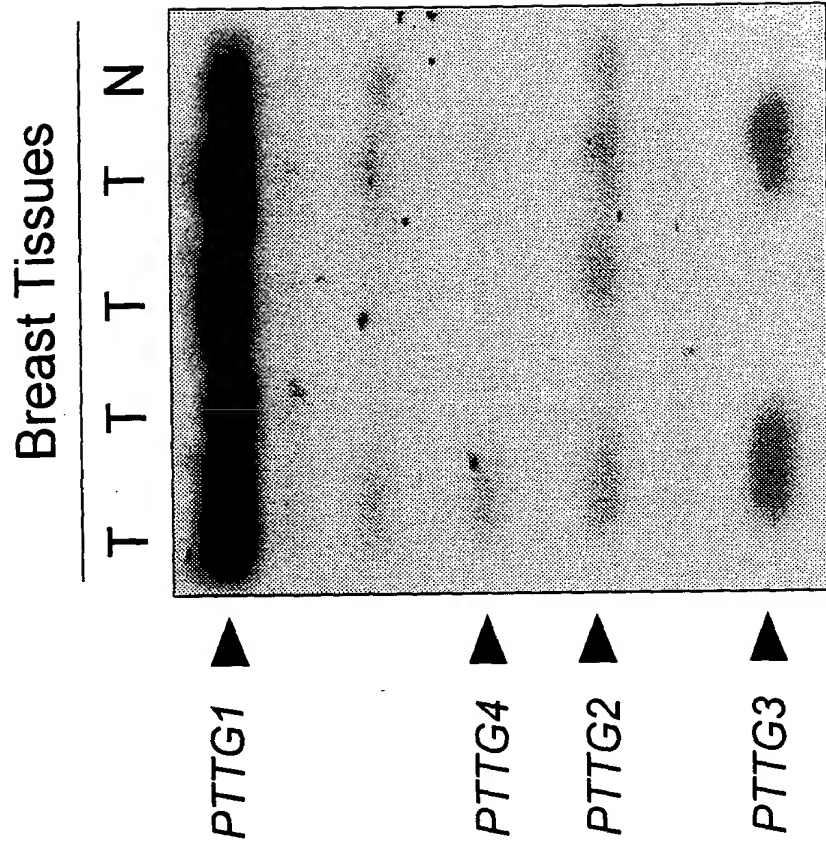


Figure 27B

PTT50" 92E45860

Ovarian Tissues

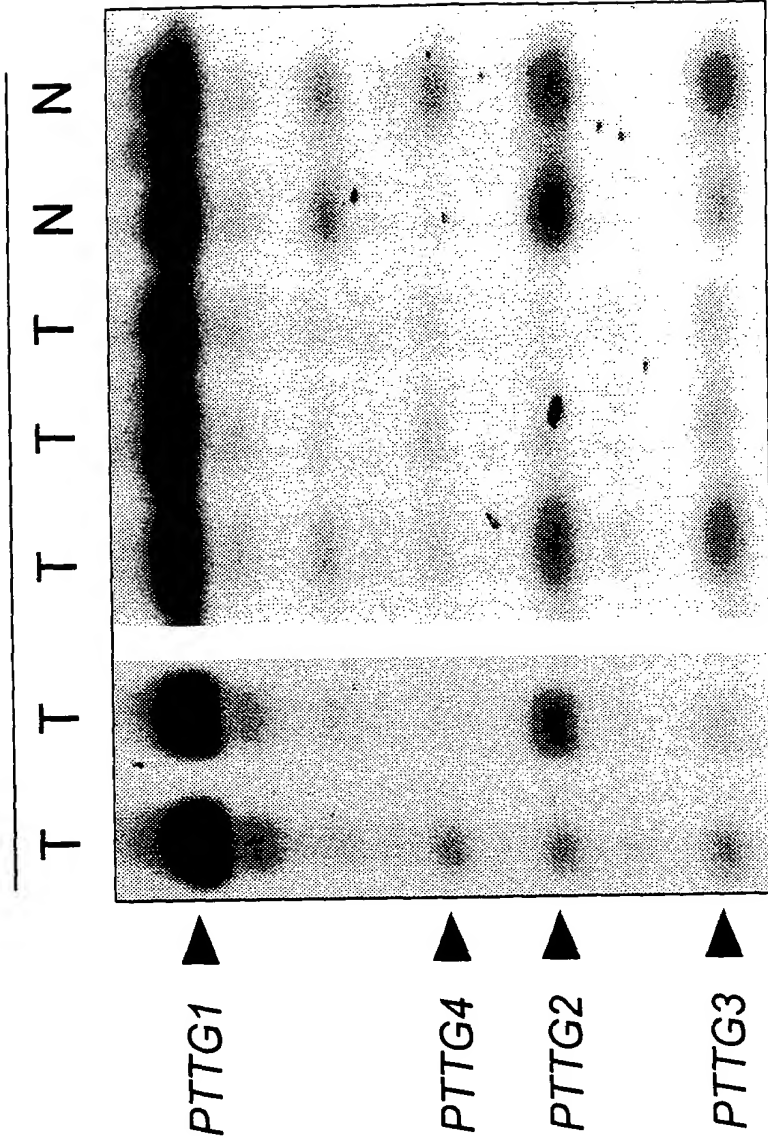


Figure 27C

09854326.051101

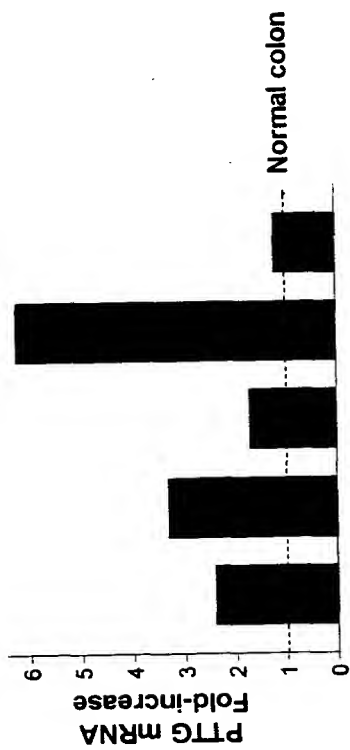


Figure 28

FOF50" 92645860

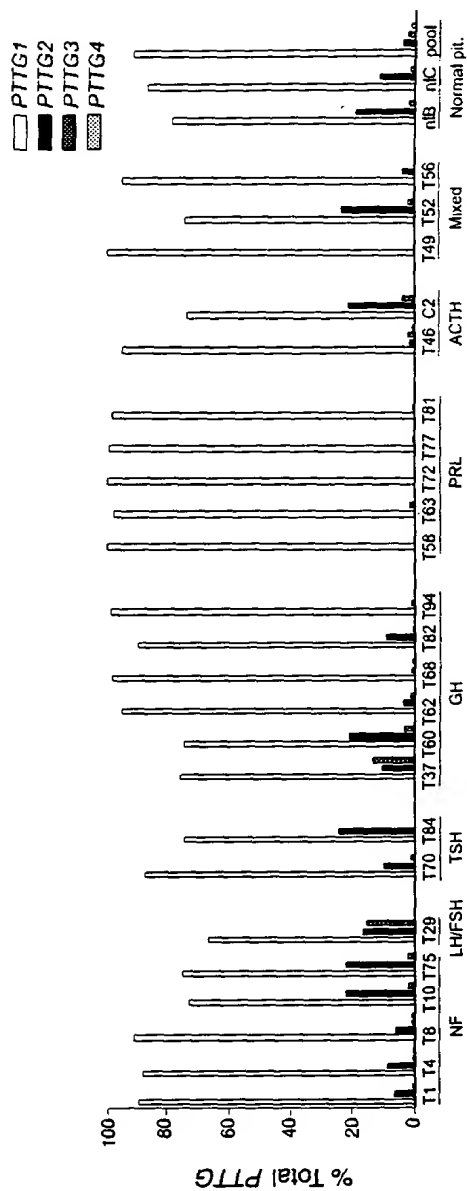


Figure 29

Interspecific hybrids

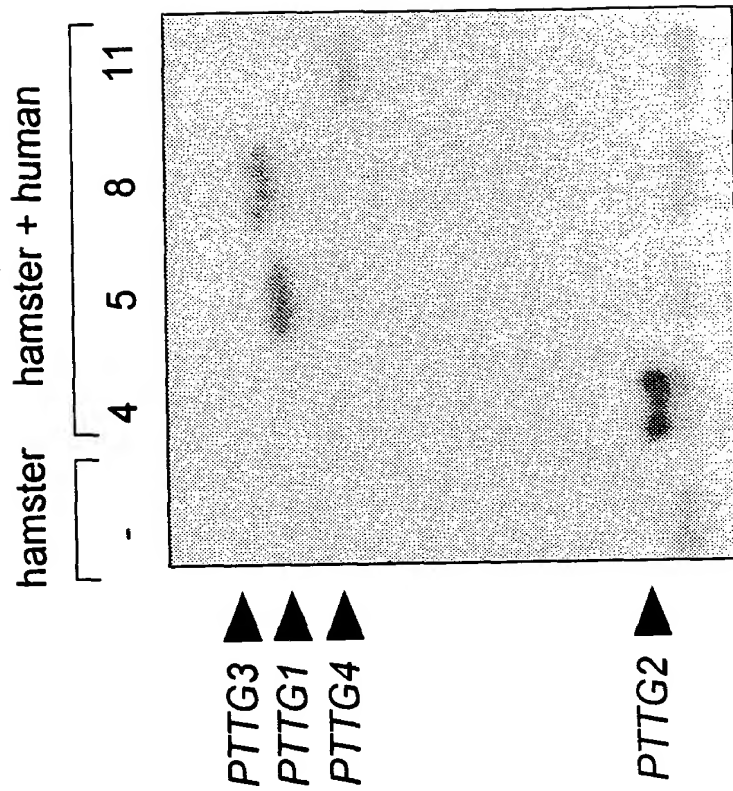


Figure 30A

FOU50" 92E45860

Colon Tissues

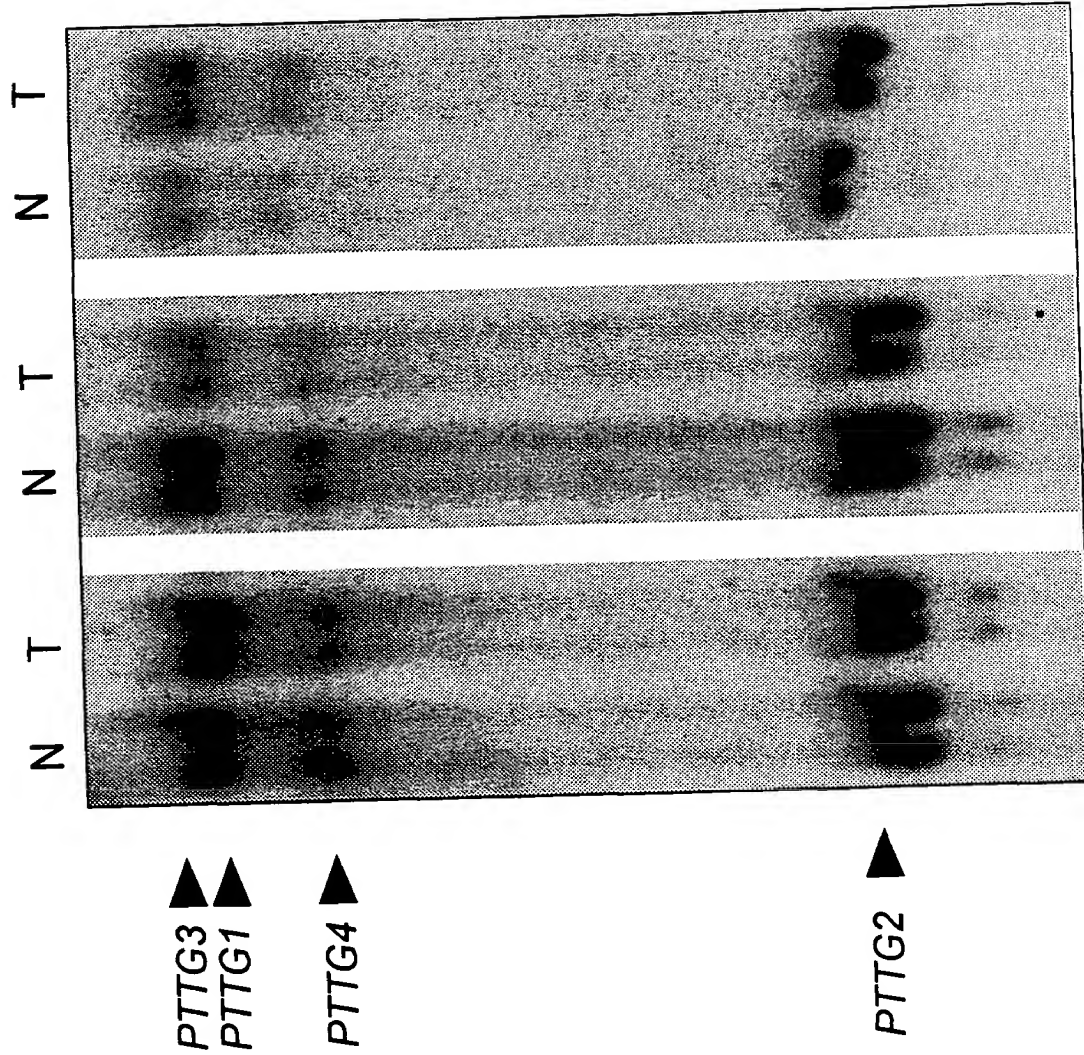


Figure 30B

09854326.051101

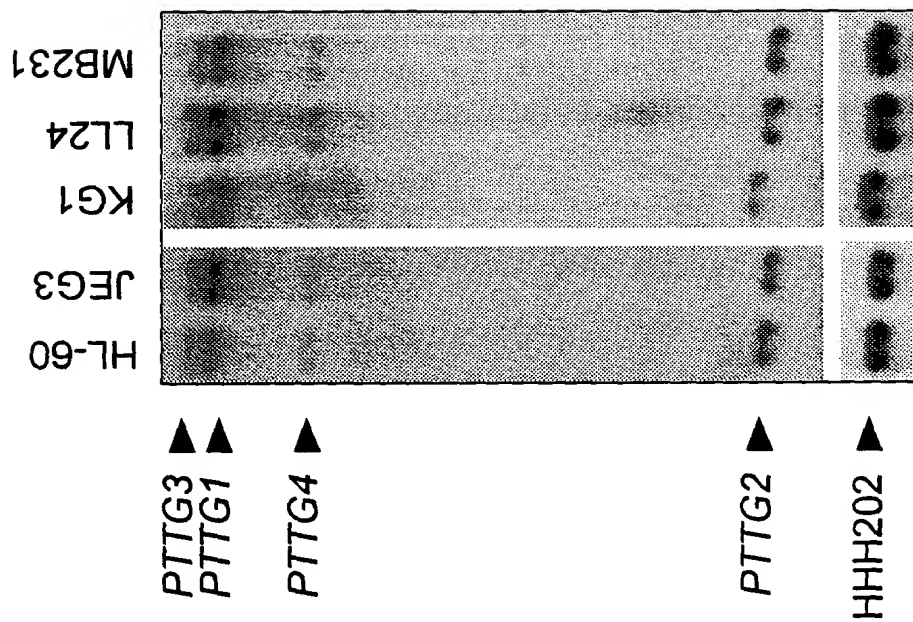


Figure 30C

09854325.051101

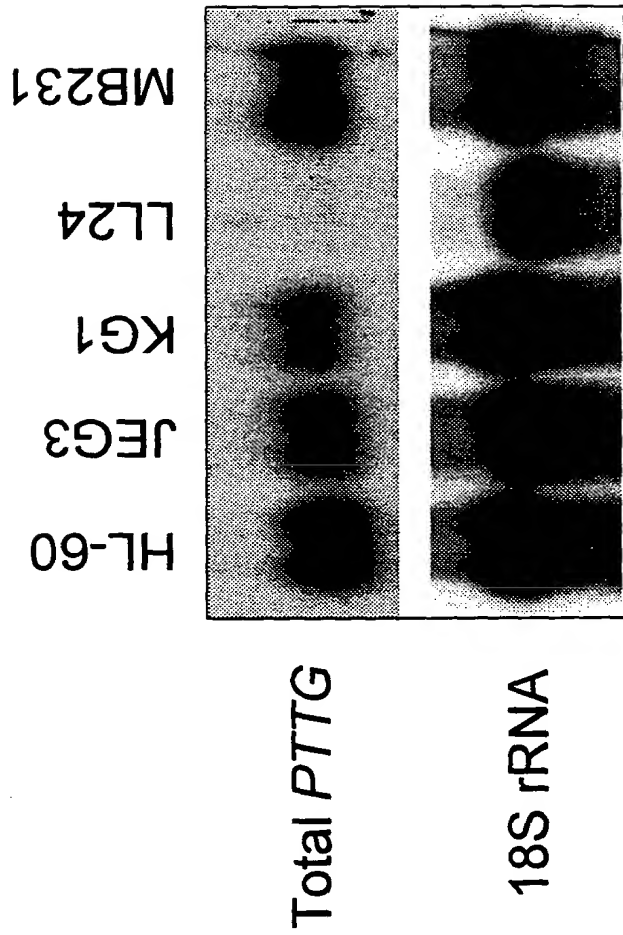


Figure 30A

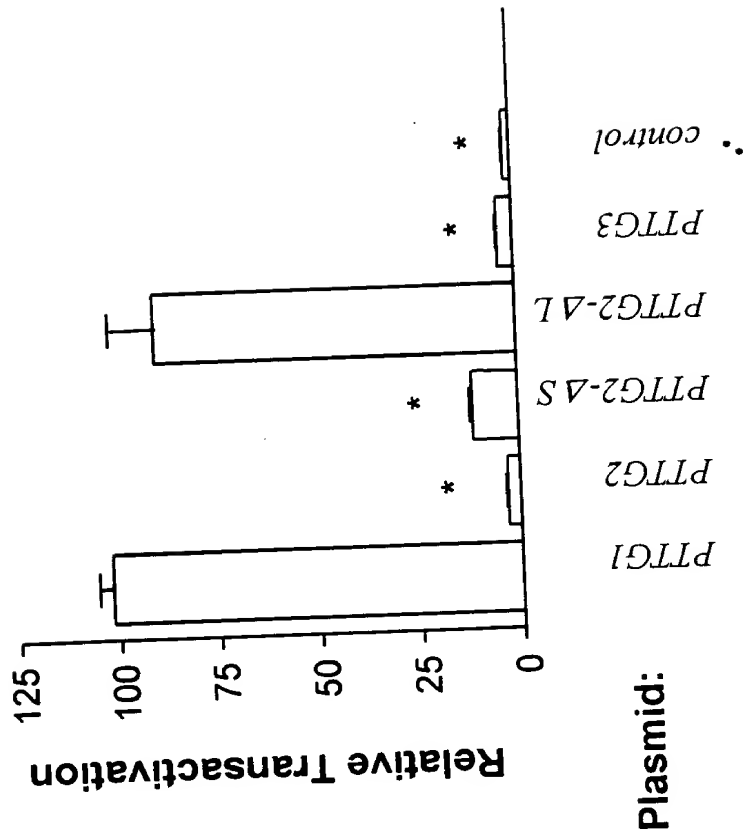


Figure 31

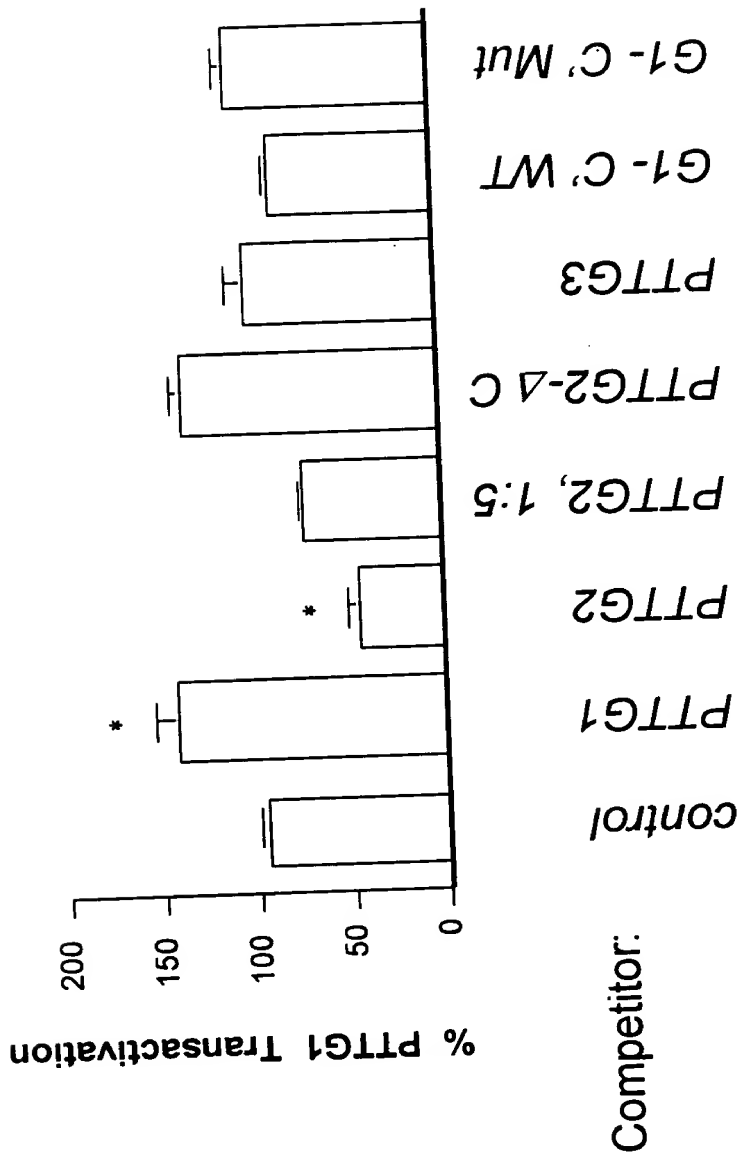


Figure 32A

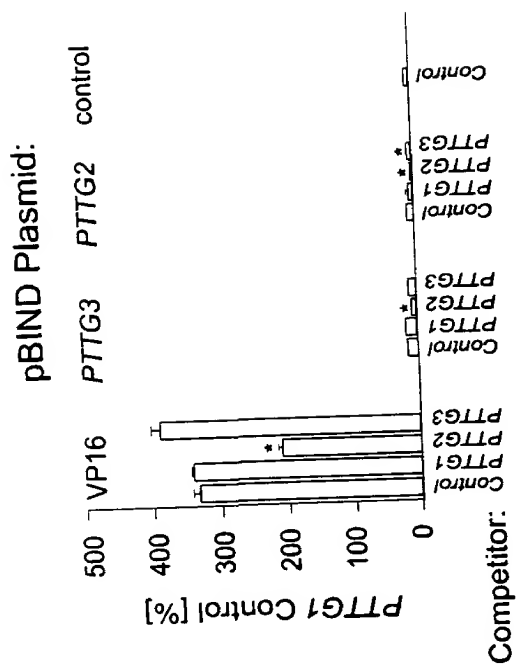


Figure 32B